

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/526, S21
PST
3/11/05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses.

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - cPAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

BEST AVAILABLE COPY

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/5261521

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS WHICH WERE INSERTED BY PTO SOFTWARE

1 Wrapped Nucleics
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 Misaligned Amino Numbering The numbering under each 3rd amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 PatentIn 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s). Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 Skipped Sequences (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO X (insert SEQ ID NO where "X" is shown)
(1) SEQUENCE CHARACTERISTICS (Do not insert any subheadings under this heading)
(11) SEQUENCE DESCRIPTION SEQ ID NO X (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(11) NUMBER OF SEQUENCES" response to include the skipped sequences.

8 Skipped Sequences (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
<210> sequence id number
<400> sequence id number
000

9 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 Invalid <211> Response Per 1.823 of Sequence Rules, the only valid <211> responses are Unknown, Artificial Sequence, or Scientific name (Genus/Species). <220>-<223> section is required when <211> response is Unknown.

11 Use of <220> Sequence(s) missing the <220> feature and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <211> "Organism" response is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (See 1.823 of Sequence Rules)

12 PatentIn 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid.



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:29

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

3 <110> APPLICANT: HECKER, MARKUS
 4 WAGNER, ANDREAS
 5 <120> TITLE OF INVENTION: Decoy-Oligonucleotide-Inhibition of CD40-Expression
 6 <130> FILE REFERENCE: DEBE:054US
 C--> 7 <140> CURRENT APPLICATION NUMBER: US/10/526,521
 C--> 9 <141> CURRENT FILING DATE: 2005-03-01
 9 <150> PRIOR APPLICATION NUMBER: PCT/DE 03/02900 <151> 2003-09-02
 W--> 10 <150> PRIOR APPLICATION NUMBER: 102 40 417.8 <151> 2002-09-02
 11 <160> NUMBER OF SEQ ID NOS: 48
 12 <170> SOFTWARE: PatentIn version 3.1
 14 <210> SEQ ID NO: 1
 15 <211> LENGTH: 9
 16 <212> TYPE: DNA
 17 <213> ORGANISM: synthetic sequence
 19 <220> FEATURE:
 20 <221> NAME/KEY: misc_feature
 21 <222> LOCATION: (1)..(1)
 22 <223> OTHER INFORMATION: V = g or c or a
 25 <220> FEATURE:
 W--> 26 <221> NAME/KEY: Decoy-Oligonucleotide
 27 <222> LOCATION: (1)..(9)
 28 <223> OTHER INFORMATION:
 31 <220> FEATURE:
 32 <221> NAME/KEY: misc_feature
 33 <222> LOCATION: (9)..(9)
 34 <223> OTHER INFORMATION: S = g or c
 W--> 37 <400> 1
 38 vtgagtcas
 41 <210> SEQ ID NO: 2
 42 <211> LENGTH: 9
 43 <212> TYPE: DNA
 44 <213> ORGANISM: synthetic sequence
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 47 <221> NAME/KEY: misc_feature
 48 <222> LOCATION: (1)..(1)
 49 <223> OTHER INFORMATION: S = g or c
 52 <220> FEATURE:
 W--> 53 <221> NAME/KEY: Decoy-Oligonucleotide
 54 <222> LOCATION: (1)..(9)
 55 <223> OTHER INFORMATION:
 58 <220> FEATURE:
 59 <221> NAME/KEY: misc_feature
 60 <222> LOCATION: (9)..(9)

move to next line

Invalid response

(pg. 1-5)

Invalid response

Mandatory
<213> response
has to be
either Artificial,
Unknown or
Genus/Species.

See item #10 on error
Summary sheet 3/11/05

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:29

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

61 <223> OTHER INFORMATION: B = g or t or c

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65 stgastcab

68 <210> SEQ ID NO: 3

69 <211> LENGTH: 21

70 <212> TYPE: DNA

71 <213> ORGANISM: synthetic sequence

73 <220> FEATURE:

W--> 74 <221> NAME/KEY: Decoy-Oligonucleotide

75 <222> LOCATION: (1)..(21)

76 <223> OTHER INFORMATION:

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80 cgcttgatga ctcagccgga a

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84 <211> LENGTH: 21

85 <212> TYPE: DNA

86 <213> ORGANISM: synthetic sequence

88 <220> FEATURE:

W--> 89 <221> NAME/KEY: Decoy-Oligonucleotide

90 <222> LOCATION: (1)..(21)

91 <223> OTHER INFORMATION:

W--> 94 <400> 4

95 ttccggtga gtcacaaagc g

98 <210> SEQ ID NO: 5

99 <211> LENGTH: 15

100 <212> TYPE: DNA

101 <213> ORGANISM: synthetic sequence

103 <220> FEATURE:

W--> 104 <221> NAME/KEY: Decoy-Oligonucleotide

105 <222> LOCATION: (1)..(15)

106 <223> OTHER INFORMATION:

W--> 109 <400> 5

110 gtgctgactc agcac

113 <210> SEQ ID NO: 6

114 <211> LENGTH: 15

115 <212> TYPE: DNA

116 <213> ORGANISM: synthetic sequence

118 <220> FEATURE:

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120 <222> LOCATION: (1)..(15)

121 <223> OTHER INFORMATION:

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125 gtgctgagtc agcac

128 <210> SEQ ID NO: 7

129 <211> LENGTH: 15

130 <212> TYPE: DNA

131 <213> ORGANISM: synthetic sequence

133 <220> FEATURE:

W--> 134 <221> NAME/KEY: Decoy-Oligonucleotide

9

See item
#10 on
error
summary
sheet.

15

15

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:29

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

✓ SAME
errors
15

135 <222> LOCATION: (1)..(15)
136 <223> OTHER INFORMATION:
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140 gtggtgactc accac
143 <210> SEQ ID NO: 8
144 <211> LENGTH: 15
145 <212> TYPE: DNA
146 <213> ORGANISM: (synthetic sequence)
148 <220> FEATURE:
W--> 149 <221> NAME/KEY: Decoy-Oligonucleotide
150 <222> LOCATION: (1)..(15)
151 <223> OTHER INFORMATION:
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155 gtggtgagtc accac 15
158 <210> SEQ ID NO: 9
159 <211> LENGTH: 17
160 <212> TYPE: DNA
161 <213> ORGANISM: (synthetic sequence)
163 <220> FEATURE:
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165 <222> LOCATION: (1)..(17)
166 <223> OTHER INFORMATION:
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175 <212> TYPE: DNA
176 <213> ORGANISM: (synthetic sequence)
178 <220> FEATURE:
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180 <222> LOCATION: (1)..(17)
181 <223> OTHER INFORMATION:
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185 agtgggtgagt caccact 17
188 <210> SEQ ID NO: 11
189 <211> LENGTH: 17
190 <212> TYPE: DNA
191 <213> ORGANISM: (synthetic sequence)
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195 <222> LOCATION: (1)..(17)
196 <223> OTHER INFORMATION:
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203 <210> SEQ ID NO: 12
204 <211> LENGTH: 17
205 <212> TYPE: DNA
206 <213> ORGANISM: (synthetic sequence)
208 <220> FEATURE:

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:29

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

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      211 <223> OTHER INFORMATION:
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      215 tgtgctgagt cagcaca 17
      218 <210> SEQ ID NO: 13
      219 <211> LENGTH: 19
      220 <212> TYPE: DNA
      221 <213> ORGANISM: synthetic sequence
      223 <220> FEATURE:
W--> 224 <221> NAME/KEY: Decoy-Oligonucleotide
      225 <222> LOCATION: (1)..(19)
      226 <223> OTHER INFORMATION:
W--> 229 <400> 13
      230 ttgtgctgac tcagcacaa 19
      233 <210> SEQ ID NO: 14
      234 <211> LENGTH: 19
      235 <212> TYPE: DNA
      236 <213> ORGANISM: synthetic sequence
      238 <220> FEATURE:
W--> 239 <221> NAME/KEY: Decoy-Oligonucleotide
      240 <222> LOCATION: (1)..(19)
      241 <223> OTHER INFORMATION:
W--> 244 <400> 14
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      249 <211> LENGTH: 13
      250 <212> TYPE: DNA
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      255 <222> LOCATION: (1)..(13)
      256 <223> OTHER INFORMATION:
W--> 259 <400> 15
      260 tggtagtca cca 13
      263 <210> SEQ ID NO: 16
      264 <211> LENGTH: 13
      265 <212> TYPE: DNA
      266 <213> ORGANISM: synthetic sequence
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      270 <222> LOCATION: (1)..(13)
      271 <223> OTHER INFORMATION:
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      275 tggtagtca cca 13
      278 <210> SEQ ID NO: 17
      279 <211> LENGTH: 15
      280 <212> TYPE: DNA
      281 <213> ORGANISM: synthetic sequence

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:29

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

283 <220> FEATURE:
W--> 284 <221> NAME/KEY: Decoy-Oligonucleotide
285 <222> LOCATION: (1)..(15)
286 <223> OTHER INFORMATION:
W--> 289 <400> 17
290 atggtgagtc accat
293 <210> SEQ ID NO: 18
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296 <213> ORGANISM: synthetic sequence
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W--> 299 <221> NAME/KEY: Decoy-Oligonucleotide
300 <222> LOCATION: (1)..(15)
301 <223> OTHER INFORMATION:
W--> 304 <400> 18
305 atggtgactc accat
308 <210> SEQ ID NO: 19
309 <211> LENGTH: 17
310 <212> TYPE: DNA
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313 <220> FEATURE:
W--> 314 <221> NAME/KEY: Decoy-Oligonucleotide
315 <222> LOCATION: (1)..(17)
316 <223> OTHER INFORMATION:
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320 tatggtgagt caccata
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324 <211> LENGTH: 17
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330 <222> LOCATION: (1)..(17)
331 <223> OTHER INFORMATION:
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335 tatggtgact caccata
338 <210> SEQ ID NO: 21
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346 <223> OTHER INFORMATION:
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350 ctatggtgag tcaccatag
353 <210> SEQ ID NO: 22
354 <211> LENGTH: 19
355 <212> TYPE: DNA

✓ SAME
15
errors

15

17

17

19

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:30

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

L:7 M:270 C: Current Application Number differs, Replaced Current Application Number
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:10 M:289 W: Identifier Missing or Out-Of-Order, <150> PRIOR APP FILING DATE
L:26 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:37 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:28
L:53 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:64 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:55
L:74 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:79 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:76
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:30

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/526,521

DATE: 03/11/2005

TIME: 08:32:30

Input Set : E:\DEBE054.txt

Output Set: N:\CRF4\03112005\J526521.raw

L:749 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:48

L:754 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:48,Line#:751